Changed a file from non-ASCII to ASCII  Changed the margins in cases where the sequence text was "wrapped" down to the aged income Edited a format error in the Current Application Data section, specifically:  Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other the number instead of using an integrated the "Number of Sequences" field. The applicant spelled out a number instead of using an integrated the spelling of a mandatory field (the headings or subheadings), specifically.  Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted extra, invalid, headings used by an applicant, specifically:  Corrected an obvious error in the response, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an orror in the Number of Sequences field, specifically:	r: 10/005, 200 Edited by:
Edited a format error in the Current Application Data section, specifically:  Edited the Current Application Data section with the actual current number. The number inputted by tapplicant was   the prior application data; or   other   Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integrated the spelling of a mandatory field (the headings or subheadings), specifically;  Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: One-ASCII "garbage" at the beginning/end of files: secretary initials/filename at end page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	
Edited the Current Application Data section with the actual current number. The number inputted by tapplicant was the prior application data; or other  Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integrated the spelling of a mandatory field (the headings or subheadings), specifically;  Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:	ed the margins in cases where the sequence text was "wrapped" down to the next time.
Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integration of the spelling of a mandatory field (the headings or subheadings), specifically.  Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: non-ASCII garbage at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	a formal error in the Current Application Data section, specifically:
Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integration of the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as  Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	the Current Application Data section with the actual current number. The number inputted by the time that the prior application data; or
Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Defeted extra, invalid, headings used by an applicant, specifically:  Defeted: non-ASCII garbage at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as  Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	the mandatory heading and subheadings for *Current Application Data*.
Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Defeted extra, invalid, headings used by an applicant, specifically:  Defeted: non-ASCII garbage at the beginning/end of files: secretary initials/filename at end page numbers throughout text; other invalid text, such as lnserted mandatory headings, specifically:  Corrected an obvious erro: in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	the "Number of Sequences" field. The applicant spelled out a number instead of using an integ
Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: non-ASCII 'garbage' at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	ed the spelling of a mandatory field (the headings or subheadings), specifically:
Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	ed the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
Deleted extra, invalid, headings used by an applicant, specifically:  Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	ed subheading placement. All responses must be on the same line as each subheading. If the nt placed a response below the subheading, this was moved to its appropriate place.
Deleted: non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	d colons after headings/subheadings. Headings edited included:
page numbers throughout text; other invalid text, such as	extra, invalid, headings used by an applicant, specifically:
Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	d: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of genumbers throughout text; other invalid text, such as
Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	d mandatory headings, specifically:
Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:	
•	
er en	
A "Hard Page Break" code was inserted by the applicant. All occurrences had to be detected.	Page Break* code was inserted by the applicant. All occurrences had to be deleted.
Deleted ending stop codon in amino acid sequences and adjusted the *(A)Length:* field accordingly (edue to a Patentin bug). Sequences corrected:	endIng stop codon in amino acid sequences and adjusted the *(A)Length:* field accordingly (e PatentIn bug). Sequences corrected:
Other:	

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

2000

OIPE

RAW SEQUENCE LISTING

DATE: 12/12/2001

PATENT APPLICATION: US/10/005,200

TIME: 14:19:50

Input Set : A:\Sequences for submission.txt Output Set: N:\CRF3\12112001\I005200.raw

**Does Not Comply** Corrected Diskette Needed

4 <110> APPLICANT: DeltaGen, Inc.

6 <120> TITLE OF INVENTION: TRANSGENIC MICE CONTAINING RPTPB

TYROSINE PHOSPHATASE GENE DISRUPTIONS

10 <130> FILE REFERENCE: 741 PCT

C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/005,200

C--> 12 <141> CURRENT FILING DATE: 2001-12-04

12 <150> PRIOR APPLICATION NUMBER: US 60/251,897

13 <151> PRIOR FILING DATE: 2000-12-06

15 <150> PRIOR APPLICATION NUMBER: US 60/302,260

16 <151> PRIOR FILING DATE: 2001-06-28

18 <160> NUMBER OF SEQ ID NOS: 3

20 <170> SOFTWARE: FastSEQ for Windows Version 4.0

## **ERRORED SEQUENCES**

73 <210> SEQ ID NO: 3

74 <211> LENGTH: 200

75 <212> TYPE: DNA

76 <213> ORGANISM: Artificial Sequence

78 <220> FEATURE:

79 <223> OTHER INFORMATION: Targeting Vector

81 <400> SEQUENCE: 3

82 ggatgagttg aagccagaac agcagcaccc tctcccttcc tacctggagt acagacacaa 60

83 egectecate egagtetace agaceaatta ttttgccage aaatgtgetg aaagteeega 120

84 cagcaqttct aaaaqtttca acattaagct tggagcagag atggacagcc tcggtggcaa 180

85 atqtqatccc aqtcaqcaqa 200

E--> 86(1

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/005,200

DATE: 12/12/2001

TIME: 14:19:51

Input Set : A:\Sequences for submission.txt
Output Set: N:\CRF3\12112001\1005200.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:86 M:254 E: No. of Bases conflict, LENGTH:Input:1 Countèd:200 SEQ:3

## OIPE

RAW SEQUENCE LISTING DATE: 12/17/2001 PATENT APPLICATION: US/10/005,200 TIME: 12:48:31

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\12172001\J005200.raw

4 <110> APPLICANT: DeltaGen, Inc. 6 <120> TITLE OF INVENTION: TRANSGENIC MICE CONTAINING RPTPB TYROSINE PHOSPHATASE GENE DISRUPTIONS 10 <130> FILE REFERENCE: 741 PCT C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/005,200 C--> 12 <141> CURRENT FILING DATE: 2001-12-04 12 <150> PRIOR APPLICATION NUMBER: US 60/251,897 13 <151> PRIOR FILING DATE: 2000-12-06 15 <150> PRIOR APPLICATION NUMBER: US 60/302,260 16 <151> PRIOR FILING DATE: 2001-06-28 18 <160> NUMBER OF SEQ ID NOS: 3 20 <170> SOFTWARE: FastSEQ for Windows Version 4.0 22 <210> SEQ ID NO: 1 23 <211> LENGTH: 1752 24 <212> TYPE: DNA 25 <213> ORGANISM: Mus musculus 27 <400> SEQUENCE: 1 28 gtcaaggaag aggtacctgg tgtccatcaa ggtgcagtcg gccggcatga ccagtgaggt 60 29 ggttgaagat agcaccatca ccatgataga ccgcccgcct caaccgcctc cacacatccg 120 30 tgtgaatgaa aaggatgtgc taatcagcaa atcttccatc aactttactg tcaactgcag 180 31 ctggttcagc gacaccaacg gagcggttgg gtactttgct gtggtggtga gagaggccga 240 32 cagcatggat gagttgaagc cagaacagca gcaccctctc ccttcctacc tggagtacag 300 33 acacaacgcc tccatccgag tctaccagac caattatttt gccagcaaat gtgctgaaag 360 34 tocogacago agttotaaaa gtttoaacat taagottgga goagagatgg acagootogg 420 35 tggcaaatgt gatcccagtc agcagaaatt ctgtgatgga ccgctgttgc cacacccgc 480 36 ctacagaatc agcatccggg cttttacaca gctatttgac gaggacttga aagagttcac 540 37 caaacctctc tactcggata cgttcttctc tatgcccatc accacagagt cagagccctt 600 38 gtttggagtt attgaaggtg tgagtgctgg cctgtttcta attggcatgc tggtggccct 660 39 tgttgccttc ttcatctgca gacagaaagc tagccacagc agggaaaggc catctgcccg 720 40 gctcagcatt cgtagggacc ggcctttgtc tgtccatctg aatctgggcc agaaaggcaa 780 41 coggaaaact tottgoocca taaagatcaa toagtttgaa gggoatttoa tgaagotgoa 840 42 ggcagactcc aactaccttc tatccaagga atatgaggac ttaaaagacg tgggtagaag 900 43 ccagtcatgt gacattgccc tcttgcctga gaatcgaggg aaaaatcgat acaacaacat 960 44 attgeettat gatgeeteaa gagtgaaget etegaatgte gatgaegaee ettgetetga 1020 45 ctacatcaac gccagctaca tccccggtaa caacttcaga cgagaataca tcgccactca 1080 46 gggaccgctt ccaggcacca aggatgactt ctggaagatg gcgtgggagc agaacgttca 1140 47 caacatcgtc atggtgaccc agtgtgttga aaagggccga gtgaagtgtg accattactg 1200 48 gccagcagac caggaccccc tctactacgg tgatctcatc ctacagatgg tctcggagtc 1260 49 cgtgctcccc gagtggacca tcagggagtt taagatatgc agtgaagaac agttggatgc 1320 50 acacagacte atecgteact tteactacae ggtgtggeea gaccatgggg teccagagae 1380 51 cacccagtcc ctgatccaat ttgtgaggac agtcagggac tacatcaaca gaagccccgg 1440 52 ggctgggccc tccgtagtgc actgcagcgc tggtgtgggc agaacaggga cgttcgttgc 1500 53 cctggaccgg atcctccagc agttggactc taaggactcc gtggacattt atggggcagt 1560 54 gcatgaccta agactccaca gggttcacat ggtccagacc gagtgtcaat atgtgtatct 1620 55 gcatcagtgt gtaagagacg tctcagagca aagaaactgc ggaaacgagc aagagaaagg 1680 56 gggtgtttcg atttatgaga atgtgaatca gagtatcaca gagatgcaat ctactcgaga 1740

57 cattaagaat tc

1752



RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/005,200

DATE: 12/17/2001 TIME: 12:48:31

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\12172001\J005200.raw

- 59 <210> SEQ ID NO: 2
- 60 <211> LENGTH: 200
- 61 <212> TYPE: DNA
- 62 <213> ORGANISM: Artificial Sequence
- 64 <220> FEATURE:
- 65 <223> OTHER INFORMATION: Targeting Vector
- 67 <400> SEQUENCE: 2
- 68 geogeoceca gaactecacg geoattgeet getettggat acctectgae teegactttg 60
- 69 atggctacag cattgagtgc cgaaaaatgg atacccaaga aatcgagttt tccagaaagc 120
- 70 tggagaaaga aaaatcactg ctcaacatca tgatgttagt acctcataag aggtacctgg 180
- 71 tgtccatcaa ggtgcagtcg
- 73 <210> SEQ ID NO: 3
- 74 <211> LENGTH: 200
- 75 <212> TYPE: DNA
- 76 <213> ORGANISM: Artificial Sequence
- 78 <220> FEATURE:
- 79 <223> OTHER INFORMATION: Targeting Vector
- 81 <400> SEQUENCE: 3
- 82 ggatgagttg aagccagaac agcagcaccc tctcccttcc tacctggagt acagacacaa 60
- 83 cgcctccatc cgagtctacc agaccaatta ttttgccagc aaatgtgctg aaagtcccga 120
- 84 cagcagttct aaaagtttca acattaagct tggagcagag atggacagcc tcggtggcaa 180
- 85 atgtgatccc agtcagcaga 20



VERIFICATION SUMMARY

PATENT APPLICATION: US/10/005,200

DATE: 12/17/2001

TIME: 12:48:32

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\12172001\J005200.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date